S. Liu

Re-run

#12

RAW SEQUENCE LISTING DATE: 09/17/2002
PATENT APPLICATION: US/09/697,863 TIME: 14:13:58

Input Set : N:\paola\US09697863A.RAW
Output Set: N:\CRF4\09172002\I697863.raw

ENTERED

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     1 <110> APPLICANT: Vlaams Interuniversitair Instituut Voor Biotechnologie VZW
     2 <120> TITLE OF INVENTION: CD-40 INTERACTING AND TRAF-INTERACTING PROTEINS
     3 <130> FILE REFERENCE: 2676-4555US
C--> 4 <140> CURRENT APPLICATION NUMBER: US/09/697,863
     5 <141> CURRENT FILING DATE: 2000-10-27
     6 <150> PRIOR APPLICATION NUMBER: PCT/EP99/03025
     7 <151> PRIOR FILING DATE: 1999-04-28
     8 <150> PRIOR APPLICATION NUMBER: EPO 98201392.2
     9 <151> PRIOR FILING DATE: 1998-04-29
    10 <160> NUMBER OF SEQ ID NOS: 6
    11 <170> SOFTWARE: PatentIn version 3.1
    13 <210> SEQ ID NO: 1
    14 <211> LENGTH: 1920
    15 <212> TYPE: DNA
    16 <213> ORGANISM: Homo sapiens
    17 <220> FEATURE:
    18 <221> NAME/KEY: misc_feature
    19 <222> LOCATION: (1627)..(1627)
    20 <223> OTHER INFORMATION: N stands for any nucleotide.
    21 <220> FEATURE:
    22 <221> NAME/KEY: CDS
    23 <222> LOCATION: (20)..(1108)
    24 <223> OTHER INFORMATION:
    25 <220> FEATURE:
    26 <221> NAME/KEY: misc_feature
    27 <222> LOCATION: (1849)..(1849)
    28 <223> OTHER INFORMATION: N stands for any nucleotide.
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    30       gtgcagaggc ggcaggaag atg gag ttg ggg agt tgc ctg gag ggc ggg agg      52
    31                          Met Glu Leu Gly Ser Cys Leu Glu Gly Gly Arg
    32                          1               5                   10
    33       gag gcg gcg gag gaa gag ggc gag cct gag gtg aaa aag cgg cga ctt      100
    34       Glu Ala Ala Glu Glu Glu Gly Glu Pro Glu Val Lys Lys Arg Arg Leu
    35                   15                  20                  25
    36       ctg tgt gtg gag ttt gcc tcg gtc gca agc tgc gat gcc gca gtg gct      148
    37       Leu Cys Val Glu Phe Ala Ser Val Ala Ser Cys Asp Ala Ala Val Ala
    38               30                  35                  40
    39       cag tgc ttc ctg gcc gag aac gac tgg gag atg gaa agg gct ctg aac      196
    40       Gln Cys Phe Leu Ala Glu Asn Asp Trp Glu Met Glu Arg Ala Leu Asn
    41           45                  50                  55
    42       tcc tac ttc gag cct ccg gtg gag gag agc gcc ttg gaa cgc cga cct      244
    43       Ser Tyr Phe Glu Pro Pro Val Glu Glu Ser Ala Leu Glu Arg Arg Pro
    44       60                  65                  70                  75
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```
45      gaa acc atc tct gag ccc aag acc tat gtt gac cta acc aat gaa gaa      292
46      Glu Thr Ile Ser Glu Pro Lys Thr Tyr Val Asp Leu Thr Asn Glu Glu
47                      80                  85                  90
48      aca act gat tcc acc act tct aaa atc agc cca tct gaa gat act cag      340
49      Thr Thr Asp Ser Thr Thr Ser Lys Ile Ser Pro Ser Glu Asp Thr Gln
50                  95                  100                 105
51      caa gaa aat ggc agc atg ttc tct ctc att acc tgg aat att gat gga      388
52      Gln Glu Asn Gly Ser Met Phe Ser Leu Ile Thr Trp Asn Ile Asp Gly
53              110                 115                 120
54      tta gat cta aac aat ctg tca gag agg gct cga ggg gtg tgt tcc tac      436
55      Leu Asp Leu Asn Asn Leu Ser Glu Arg Ala Arg Gly Val Cys Ser Tyr
56          125                 130                 135
57      tta gct ttg tac agc cca gat gtg ata ttt cta cag gaa gtt att ccc      484
58      Leu Ala Leu Tyr Ser Pro Asp Val Ile Phe Leu Gln Glu Val Ile Pro
59      140                 145                 150                 155
60      cca tat tat agc tac cta aag aag aga tca agt aat tat gag att att      532
61      Pro Tyr Tyr Ser Tyr Leu Lys Lys Arg Ser Ser Asn Tyr Glu Ile Ile
62                      160                 165                 170
63      aca ggt cat gaa gaa gga tat ttc aca gct ata atg ttg aag aaa tca      580
64      Thr Gly His Glu Glu Gly Tyr Phe Thr Ala Ile Met Leu Lys Lys Ser
65                  175                 180                 185
66      aga gtg aaa tta aaa agc caa gag att att cct ttt cca agt acc aaa      628
67      Arg Val Lys Leu Lys Ser Gln Glu Ile Ile Pro Phe Pro Ser Thr Lys
68              190                 195                 200
69      atg atg aga aac ctt tta tgt gtg cat gtg aat gtg tca gga aat gag      676
70      Met Met Arg Asn Leu Leu Cys Val His Val Asn Val Ser Gly Asn Glu
71          205                 210                 215
72      ctt tgc ctt atg aca tcc cat ttg gag agc acc aga ggg cat gct gcg      724
73      Leu Cys Leu Met Thr Ser His Leu Glu Ser Thr Arg Gly His Ala Ala
74      220                 225                 230                 235
75      gaa cga atg aat cag tta aaa atg gtt tta aag aaa atg caa gag gct      772
76      Glu Arg Met Asn Gln Leu Lys Met Val Leu Lys Lys Met Gln Glu Ala
77                      240                 245                 250
78      cca gag tca gct aca gtt ata ttt gca gga gat aca aat cta agg gat      820
79      Pro Glu Ser Ala Thr Val Ile Phe Ala Gly Asp Thr Asn Leu Arg Asp
80                  255                 260                 265
81      cga gag gtt acc aga tgt ggt ggt tta ccc aac aac att gtg gat gtc      868
82      Arg Glu Val Thr Arg Cys Gly Gly Leu Pro Asn Asn Ile Val Asp Val
83              270                 275                 280
84      tgg gag ttt ttg ggc aaa cct aaa cat tgc cag tat aca tgg gat aca      916
85      Trp Glu Phe Leu Gly Lys Pro Lys His Cys Gln Tyr Thr Trp Asp Thr
86          285                 290                 295
87      caa atg aac tct aat ctt gga ata act gct gct tgt aaa ctt cgt ttt      964
88      Gln Met Asn Ser Asn Leu Gly Ile Thr Ala Ala Cys Lys Leu Arg Phe
89      300                 305                 310                 315
90      gat cga ata ttt ttc aga gca gca gca gaa gag gga cac att att ccc     1012
91      Asp Arg Ile Phe Phe Arg Ala Ala Ala Glu Glu Gly His Ile Ile Pro
92                      320                 325                 330
93      cga agt ttg gac ctt ctt gga tta gaa aaa ctg gac tgt ggt aga ttt     1060
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```
Arg Ser Leu Asp Leu Leu Gly Leu Glu Lys Leu Asp Cys Gly Arg Phe
            335                 340                 345
cct agt gat cac tgg ggt ctt ctg tgc aac tta gat ata ata ttg taa     1108
Pro Ser Asp His Trp Gly Leu Leu Cys Asn Leu Asp Ile Ile Leu
        350                 355                 360
aatgcttttc aagtgtgggt tttgccctga ttgttgcaaa tacaatttcc accttctgga  1168
 aaggtaggtt tgctgtggag gaaataatgt actagatcat tgtcacagaa aaaccaacta   1228
 tgatttatgg ttgtgttttc agaattcaac attaaagatt aatgtttatt taaacgaaca   1288
 cattcctgca ttcaggatgt gaggccattt aataaaaagg gcacaaagcc tgtcagagtt   1348
 ttcaacggtg cttacagctg ccagctggat tccaaacagg taccccattg tctctgagct   1408
 aatgtttata tttttccatt caggcaccga aatagttaat atttaaaata agtcttcaaa   1468
 agaaaacata agagattatt gagttcttgg gactggatcc tttatttcat aagttcagat   1528
 catcttaaat gaaaatgcca tgattatctg cagttaagta gatgacagct attctacatc   1588
W--> agacttgatt tttgtcagct aattacataa ttggtaagnt ataattgaaa ccttatggct   1648
 taaaattcct taactccttt ttgattcatg tttgtagtca tgttgtcaac agaggcaaag   1708
 ttaagcttga tgatggttaa aatcggtttg atagcaccat gggacatttt tttaacaaaa   1768
 ataaatgcat gaagagacat agccttttag ttttgctaat tgtgaaatgg aaatgcttta   1828
W--> caggaagtaa atgcaaatta nttttaagtg tgctttaaag aaaaatattt tccccacagg   1888
 agaaatttaa ataaagaatt ttatttggta aa                                1920
<210> SEQ ID NO: 2
<211> LENGTH: 362
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
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1               5                   10                  15
Glu Gly Glu Pro Glu Val Lys Lys Arg Arg Leu Leu Cys Val Glu Phe
            20                  25                  30
Ala Ser Val Ala Ser Cys Asp Ala Ala Val Ala Gln Cys Phe Leu Ala
        35                  40                  45
Glu Asn Asp Trp Glu Met Glu Arg Ala Leu Asn Ser Tyr Phe Glu Pro
    50                  55                  60
Pro Val Glu Glu Ser Ala Leu Glu Arg Arg Pro Glu Thr Ile Ser Glu
65                  70                  75                  80
Pro Lys Thr Tyr Val Asp Leu Thr Asn Glu Glu Thr Thr Asp Ser Thr
                85                  90                  95
Thr Ser Lys Ile Ser Pro Ser Glu Asp Thr Gln Gln Glu Asn Gly Ser
            100                 105                 110
Met Phe Ser Leu Ile Thr Trp Asn Ile Asp Gly Leu Asp Leu Asn Asn
        115                 120                 125
Leu Ser Glu Arg Ala Arg Gly Val Cys Ser Tyr Leu Ala Leu Tyr Ser
    130                 135                 140
Pro Asp Val Ile Phe Leu Gln Glu Val Ile Pro Pro Tyr Tyr Ser Tyr
145                 150                 155                 160
Leu Lys Lys Arg Ser Ser Asn Tyr Glu Ile Ile Thr Gly His Glu Glu
                165                 170                 175
Gly Tyr Phe Thr Ala Ile Met Leu Lys Lys Ser Arg Val Lys Leu Lys
            180                 185                 190
Ser Gln Glu Ile Ile Pro Phe Pro Ser Thr Lys Met Met Arg Asn Leu
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RAW SEQUENCE LISTING DATE: 09/17/2002
PATENT APPLICATION: US/09/697,863 TIME: 14:13:58

Input Set : N:\paola\US09697863A.RAW
Output Set: N:\CRF4\09172002\I697863.raw

```
                195                 200                 205
Leu Cys Val His Val Asn Val Ser Gly Asn Glu Leu Cys Leu Met Thr
    210                 215                 220
Ser His Leu Glu Ser Thr Arg Gly His Ala Ala Glu Arg Met Asn Gln
225                 230                 235                 240
Leu Lys Met Val Leu Lys Lys Met Gln Glu Ala Pro Glu Ser Ala Thr
                245                 250                 255
Val Ile Phe Ala Gly Asp Thr Asn Leu Arg Asp Arg Glu Val Thr Arg
            260                 265                 270
Cys Gly Gly Leu Pro Asn Asn Ile Val Asp Val Trp Glu Phe Leu Gly
        275                 280                 285
Lys Pro Lys His Cys Gln Tyr Thr Trp Asp Thr Gln Met Asn Ser Asn
    290                 295                 300
Leu Gly Ile Thr Ala Ala Cys Lys Leu Arg Phe Asp Arg Ile Phe Phe
305                 310                 315                 320
Arg Ala Ala Ala Glu Glu Gly His Ile Ile Pro Arg Ser Leu Asp Leu
                325                 330                 335
Leu Gly Leu Glu Lys Leu Asp Cys Gly Arg Phe Pro Ser Asp His Trp
            340                 345                 350
Gly Leu Leu Cys Asn Leu Asp Ile Ile Leu
        355                 360
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<LENGTH: 1312
<TYPE: DNA
<ORGANISM: Mus musculus
<FEATURE:
<NAME/KEY: CDS
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<OTHER INFORMATION:
<SEQUENCE: 3

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aattcggcac gagggcggga agcagcgtga agagcgggtg ttttgagggg accctgcggc     120
g atg gcg tct ggc agc agt tcc gat gcg gcg gag ccc gca ggg ccg gca     169
  Met Ala Ser Gly Ser Ser Ser Asp Ala Ala Glu Pro Ala Gly Pro Ala
  1               5                  10                  15
ggg cgg gcg gcg tcg gcg ccc gaa gca gca cag gcg gag gag gac cgg       217
Gly Arg Ala Ala Ser Ala Pro Glu Ala Ala Gln Ala Glu Glu Asp Arg
              20                  25                  30
gtg aag agg cgg cgg ctt cag tgc ctg ggc ttt gcg ttg gtg ggg gga       265
Val Lys Arg Arg Arg Leu Gln Cys Leu Gly Phe Ala Leu Val Gly Gly
          35                  40                  45
tgc gac ccc acg atg gtc ccc agc gtc ctg cgg gag aac gac tgg cag       313
Cys Asp Pro Thr Met Val Pro Ser Val Leu Arg Glu Asn Asp Trp Gln
      50                  55                  60
acg cag aaa gcc ctg agc gcc tac ttc gag ctg cca gag aac gac caa       361
Thr Gln Lys Ala Leu Ser Ala Tyr Phe Glu Leu Pro Glu Asn Asp Gln
65                  70                  75                  80
ggg tgg ccg cgc cag cct ccc acg tcc ttc aag tcc gag gcc tat gtt       409
Gly Trp Pro Arg Gln Pro Pro Thr Ser Phe Lys Ser Glu Ala Tyr Val
```

RAW SEQUENCE LISTING DATE: 09/17/2002
PATENT APPLICATION: US/09/697,863 TIME: 14:13:58

Input Set : N:\paola\US09697863A.RAW
Output Set: N:\CRF4\09172002\I697863.raw

```
                    85                  90                  95
gat cta acc aac gag gat gca aat gat aca acc att tta gaa gcc agt      457
Asp Leu Thr Asn Glu Asp Ala Asn Asp Thr Thr Ile Leu Glu Ala Ser
            100                 105                 110
cca tct gga act cct cta gaa gat agc agc act att tct ttc att acc      505
Pro Ser Gly Thr Pro Leu Glu Asp Ser Ser Thr Ile Ser Phe Ile Thr
        115                 120                 125
tgg aat att gat gga tta gat gga tgc aat ctg ccc gag agg gct cga      553
Trp Asn Ile Asp Gly Leu Asp Gly Cys Asn Leu Pro Glu Arg Ala Arg
    130                 135                 140
ggg gtg tgt tcc tgc cta gct ttg tat agt cca gat gtg gta ttt cta      601
Gly Val Cys Ser Cys Leu Ala Leu Tyr Ser Pro Asp Val Val Phe Leu
145                 150                 155                 160
cag gaa gtt atc ccc cca tac tgt gcc tac cta aag aag aga gca gcc      649
Gln Glu Val Ile Pro Pro Tyr Cys Ala Tyr Leu Lys Lys Arg Ala Ala
                165                 170                 175
agt tac aca att att aca ggt aat gaa gaa gga tat ttc aca gct ata      697
Ser Tyr Thr Ile Ile Thr Gly Asn Glu Glu Gly Tyr Phe Thr Ala Ile
            180                 185                 190
cta ttg aag aaa gga aga gtg aaa ttt aaa agt cag gag att att cct      745
Leu Leu Lys Lys Gly Arg Val Lys Phe Lys Ser Gln Glu Ile Ile Pro
        195                 200                 205
ttt cca aat acc aaa atg atg aga aac ctg cta tgc gta aat gtg agt      793
Phe Pro Asn Thr Lys Met Met Arg Asn Leu Leu Cys Val Asn Val Ser
    210                 215                 220
ttg ggt gga aat gaa ttt tgc ctt atg aca tcc cat ttg gag agc acc      841
Leu Gly Gly Asn Glu Phe Cys Leu Met Thr Ser His Leu Glu Ser Thr
225                 230                 235                 240
aga gaa cat tct gcg gaa cga ata aga caa tta aaa act gtt ctt gga      889
Arg Glu His Ser Ala Glu Arg Ile Arg Gln Leu Lys Thr Val Leu Gly
                245                 250                 255
aaa atg caa gag gct cca gat tca acc acg gtt ata ttt gca gga gat      937
Lys Met Gln Glu Ala Pro Asp Ser Thr Thr Val Ile Phe Ala Gly Asp
            260                 265                 270
aca aat tta aga gat caa gaa gtt atc aaa tgt ggt ggt tta cct gac      985
Thr Asn Leu Arg Asp Gln Glu Val Ile Lys Cys Gly Gly Leu Pro Asp
        275                 280                 285
aac gtt ttt gat gcc tgg gaa ttt tta ggc aaa cct aaa cat tgc cag     1033
Asn Val Phe Asp Ala Trp Glu Phe Leu Gly Lys Pro Lys His Cys Gln
    290                 295                 300
tat aca tgg gat acg aaa gca aat aac aac ctc agg atc cct gct gct     1081
Tyr Thr Trp Asp Thr Lys Ala Asn Asn Asn Leu Arg Ile Pro Ala Ala
305                 310                 315                 320
tat aag cat cgt ttt gat cga ata ttt ttc aga gca gaa gag ggg cac     1129
Tyr Lys His Arg Phe Asp Arg Ile Phe Phe Arg Ala Glu Glu Gly His
                325                 330                 335
ctt att cct caa agt tta gac ctt gtt ggg ttg gaa aaa ctg gac tgt     1177
Leu Ile Pro Gln Ser Leu Asp Leu Val Gly Leu Glu Lys Leu Asp Cys
            340                 345                 350
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/17/2002
PATENT APPLICATION: US/09/697,863 TIME: 14:13:59

Input Set : N:\paola\US09697863A.RAW
Output Set: N:\CRF4\09172002\I697863.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 1627,1849
Seq#:5; N Pos. 1392,1530,1531
Seq#:5; Xaa Pos. 395,441
Seq#:6; Xaa Pos. 395,441

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:6; Line(s) 419

VERIFICATION SUMMARY DATE: 09/17/2002
PATENT APPLICATION: US/09/697,863 TIME: 14:13:59

Input Set : N:\paola\US09697863A.RAW
Output Set: N:\CRF4\09172002\I697863.raw

L:4 M:270 C: Current Application Number differs, Wrong Format
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1588
L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1828
L:400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:1384
L:401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:1432
L:409 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:1528
L:410 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:1536
L:476 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:384
L:482 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:432